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Figure 1A
1/14

1 ATGAGCTCCCGCATCGCCAGGGCGCTCGCCTTAGTCGTCACCCTTCTCCACTTGACCAGG 60
1 M S S R I A R A L A L V V T L L H L T R 20

61 CTGGCGCTCTCCACCTGCCCGCTGCCTGCCACTGCCCCCTGGAGGCGCCCAAGTGC GCG 120
21 L A L S T C P A A C H C P L E A P K C A 40

121 CCGGGAGTCGGGCTGGTCCGGGACGGCTGCGGCTGCTGTAAGGTCTGCGCCAAGCAGCTC 180
41 P G V G L V R D G C G C C K V C A K Q L 60

181 AACGAGGACTGCAGCAAAACGCAGCCCTGCGACCACACCAAGGGGCTGGAATGCAACTTC 240
61 N E D C S K T Q P C D H T K G L E C N F 80

241 GGC GCCAGCTCCACCGCTCTGAAGGGGATCTGCAGAGCTCAGTCAGAGGGCAGACCCTGT 300
81 G A S S T A L K G I C R A Q S E G R P C 100

301 GAATATAACTCCAGAATCTACCAAAACGGGGAAAGTTTCCAGCCCAACTGTAAACATCAG 360
101 E Y N S R I Y Q N G E S F Q P N C K H Q 120

361 TGCACATGTATTGATGGCGCCGTGGGCTGCATTCTCTGTGTCCCAAGAACTATCTCTC 420
121 C T C I D G A V G C I P L C P Q E L S L 140

421 CCCAACTTGGGCTGTCCCAACCCTCGGCTGGTCAAAGTTACCGGGCAGTGCTGCGAGGAG 480
141 P N L G C P N P R L V K V T G Q C C E E 160

481 TGGGTCTGTGACGAGGATAGTATCAAGGACCCCATGGAGGACCAGGACGGCCTCCTTGGC 540
161 W V C D E D S I K D P M E D Q D G L L G 180

541 AAGGAGCTGGGATTTCGATGCCTCCGAGGTGGAGTTGACGAGAAACAATGAATTGATTGCA 600
181 K E L G F D A S E V E L T R N N E L I A 200

601 GTTGGAAGAGGCAGCTCACTGAAGCGGCTCCCTGTTTTTGAATGGAGCCTCGCATCCTA 660
201 V G K G S S L K R L P V F G M E P R I L 220

661 TACAACCCCTTTACAAGGCCAGAAATGTATTGTTCAAACAACCTTCATGGTCCCAGTGCTCA 720
221 Y N P L Q G Q K C I V Q T T S W S Q C S 240

721 AAGACCTGTGGAACCTGGTATCTCCACACGAGTTACCAATGACAACCCTGAGTGCCGCTT 780
241 K T C G T G I S T R V T N D N P E C R L 260

781 GTGAAAGAAACCCGATTGTGTGAGGTGCGGCCCTTGTGGACAGCCAGTGTAACAGCAGCCTG 840
261 V K E T R I C E V R P C G Q P V Y S S L 280

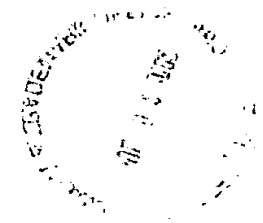


Figure 1B

2/14

841	AAAAAGGGCAAGAAATGCAGCAAGACCAAGAAATCCCCCGAACCAGTCAGGTTTACTTAC	900
281	K K G K K C S K T K K S P E P V R F T Y	300
901	GCTGGATGTTTGAGTGTGAAGAAATACCGGCCCAAGTACTGCGGTTCTGCGTGGACGGC	960
301	A G C L S V K K Y R P K Y C G S C V D G	320
961	CGATGCTGCACGCCCCAGCTGACCAGGACTGTGAAGATGCGGTTCCGCTGCGAAGATGGG	1020
321	R C C T P Q L T R T V K M R F R C E D G	340
1021	GAGACATTTTCCAAGAACGTCATGATGATCCAGTCCTGCAAATGCAACTACAACCTGCCCCG	1080
341	E T F S K N V M M I Q S C K C N Y N C P	360
1081	CATGCCAATGAAGCAGCGTTTCCCTTCTACAGGCTGTTCAATGACATTCAAAATTTAGG	1140
361	H A N E A A F P F Y R L F N D I H K F R	380
1141	GAATAA	1146
381	D *	382

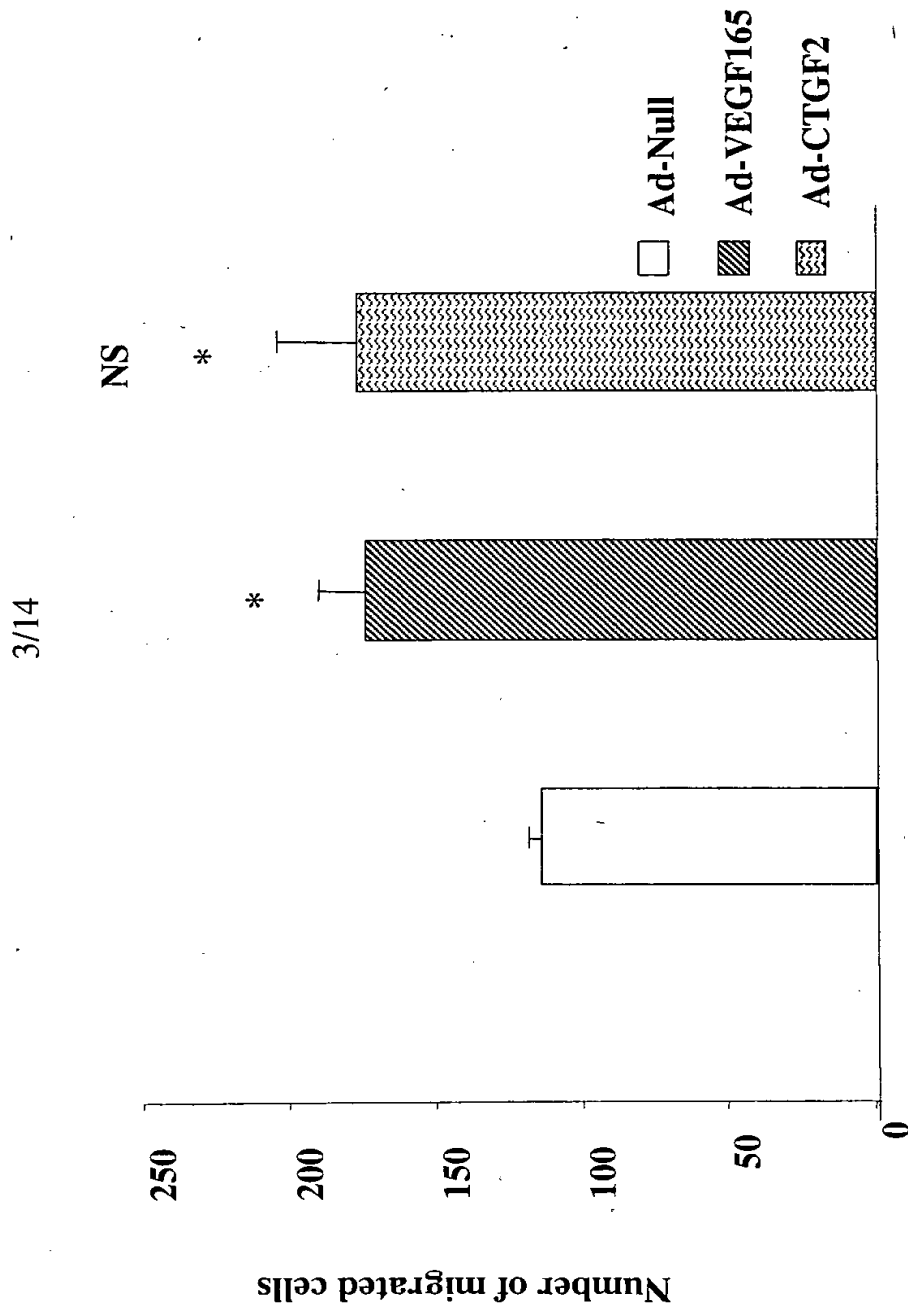
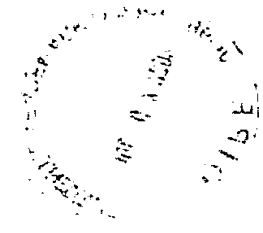
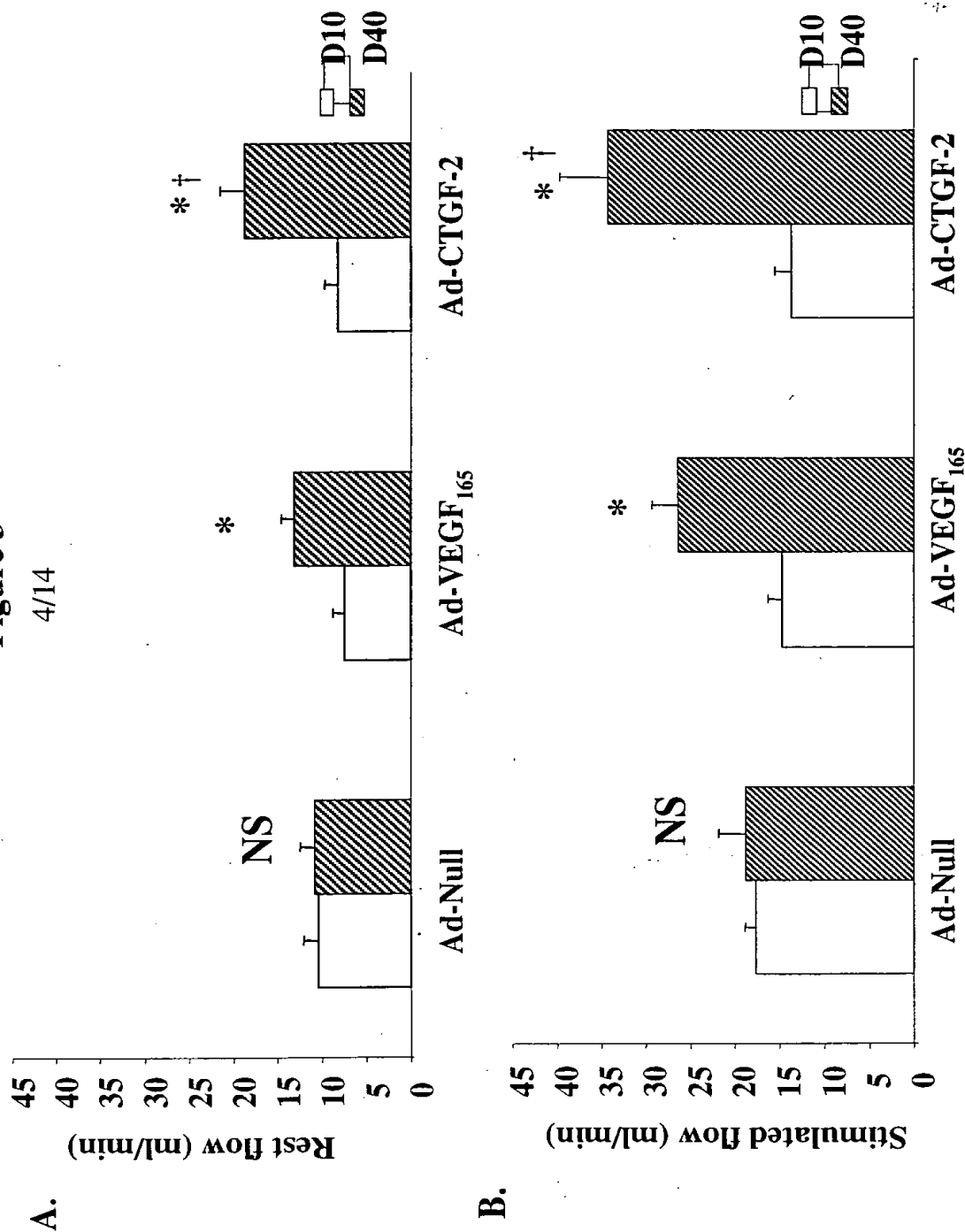


Figure 2

Figure 3

4/14



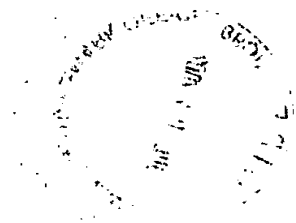
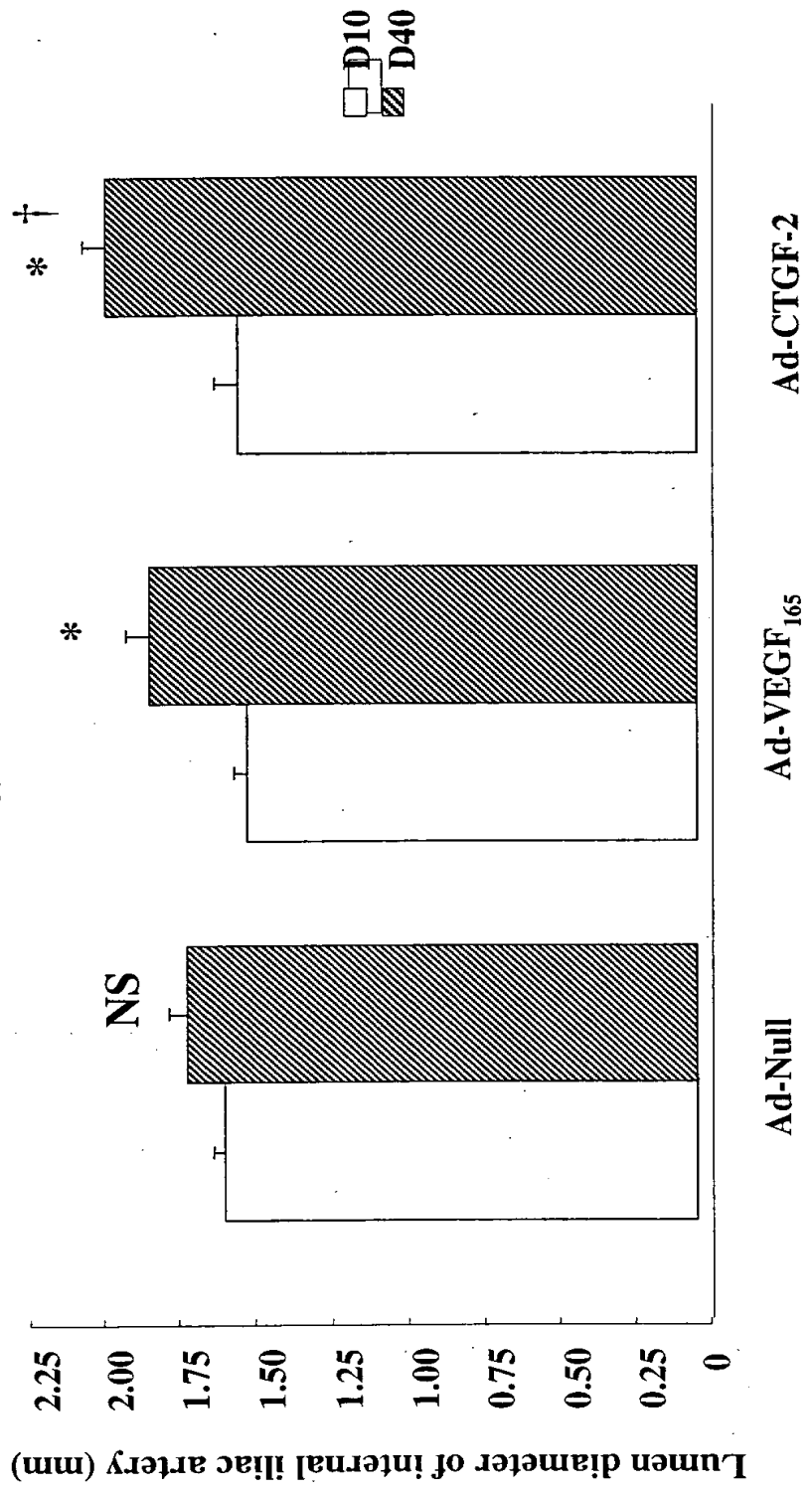


Figure 4

5/14



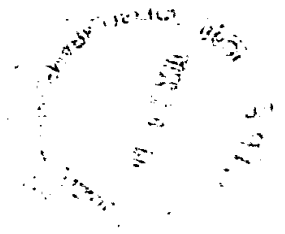
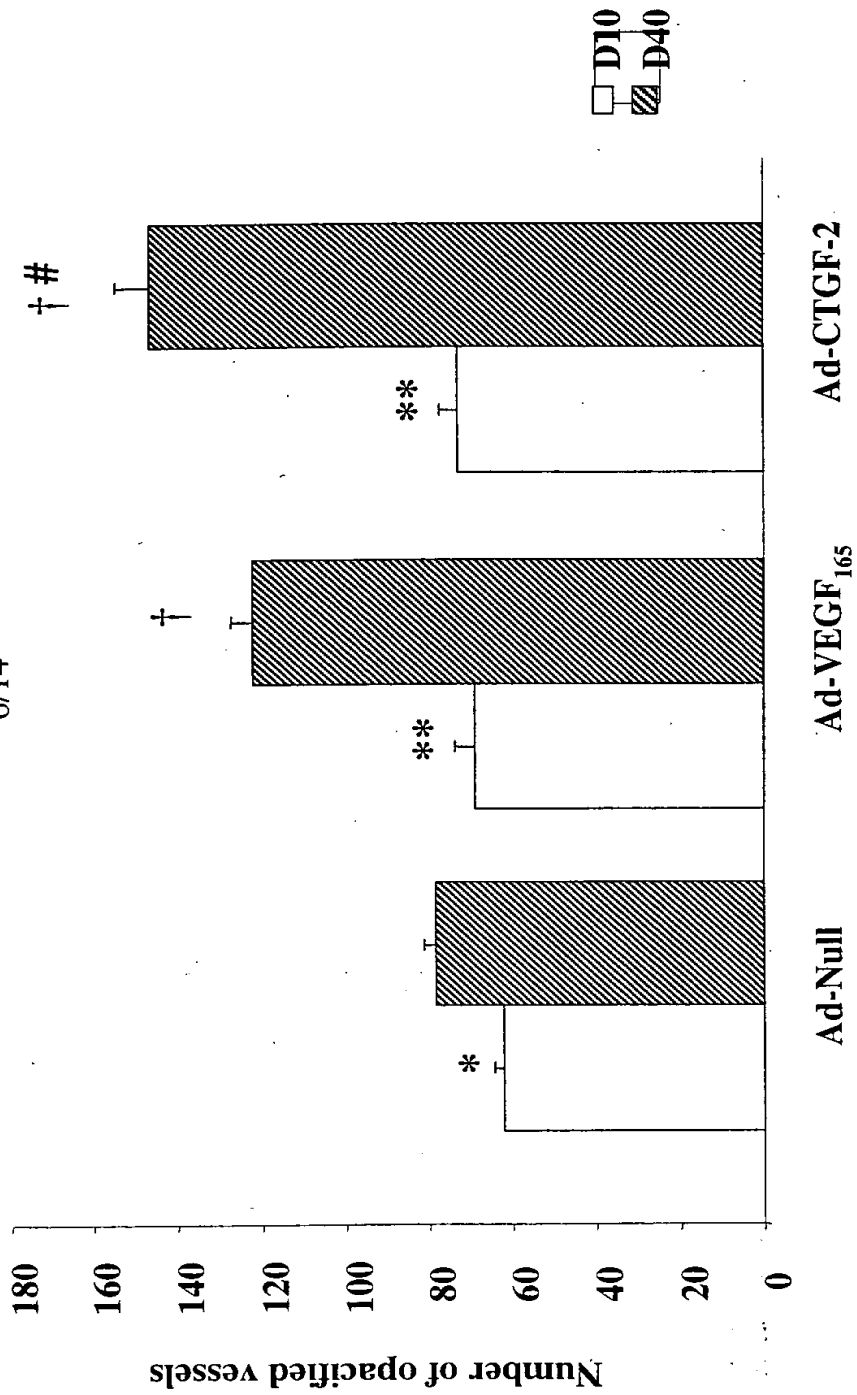


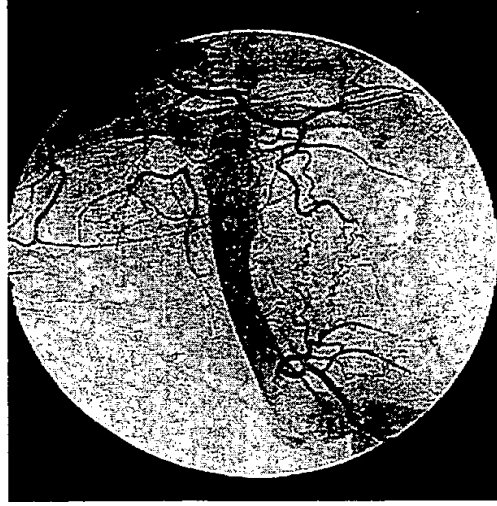
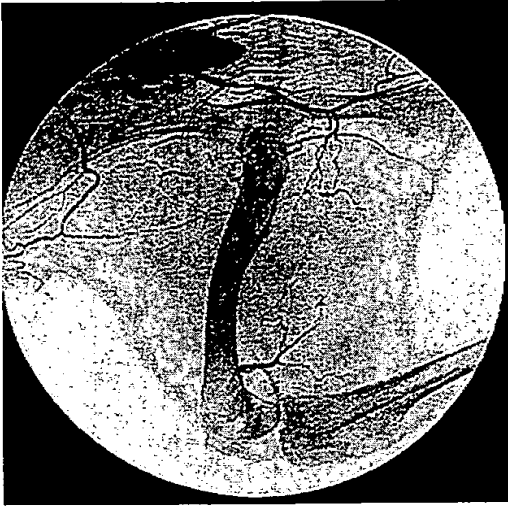
Figure 5

6/14

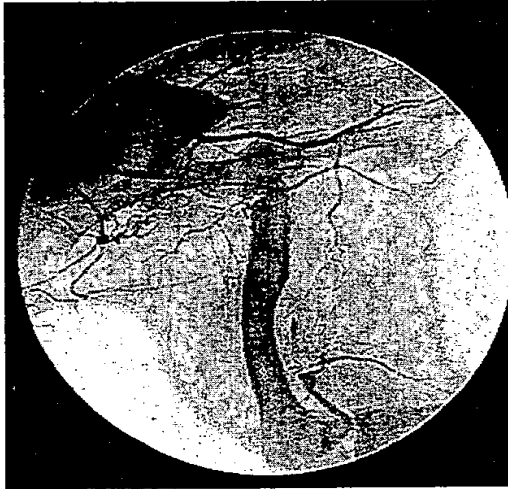


7/14

Ad-CTGF2



Ad-VEGF165



Ad-Null

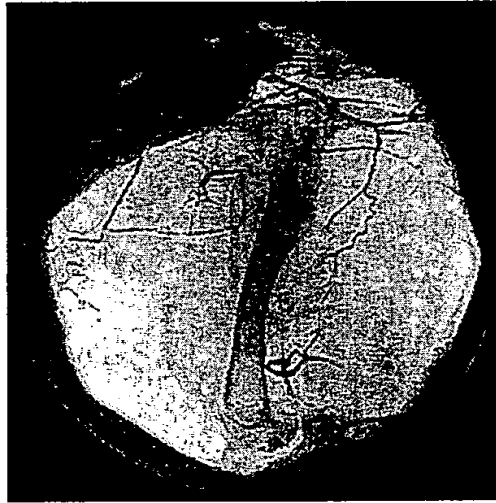
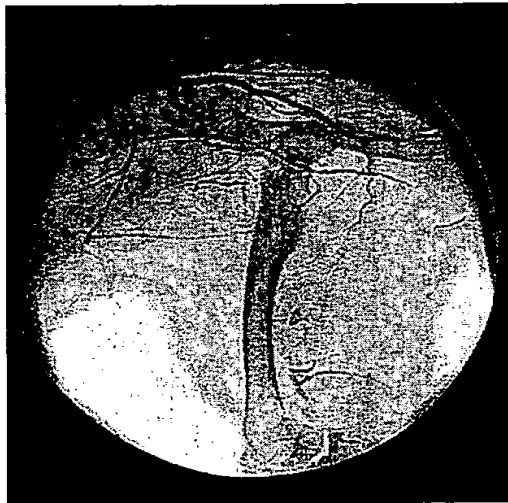
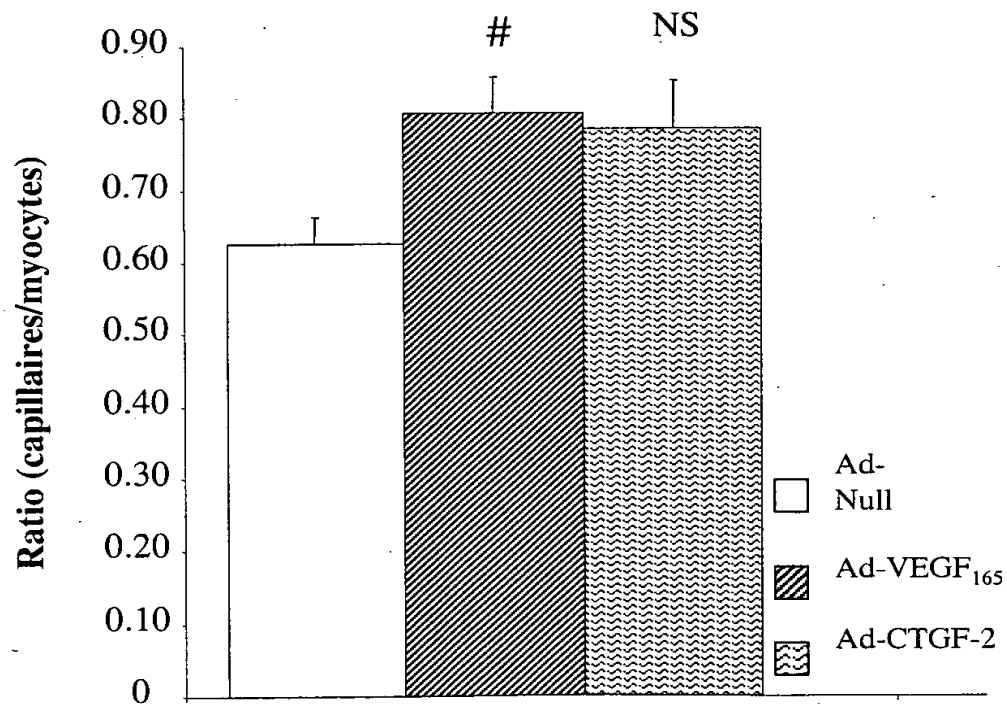
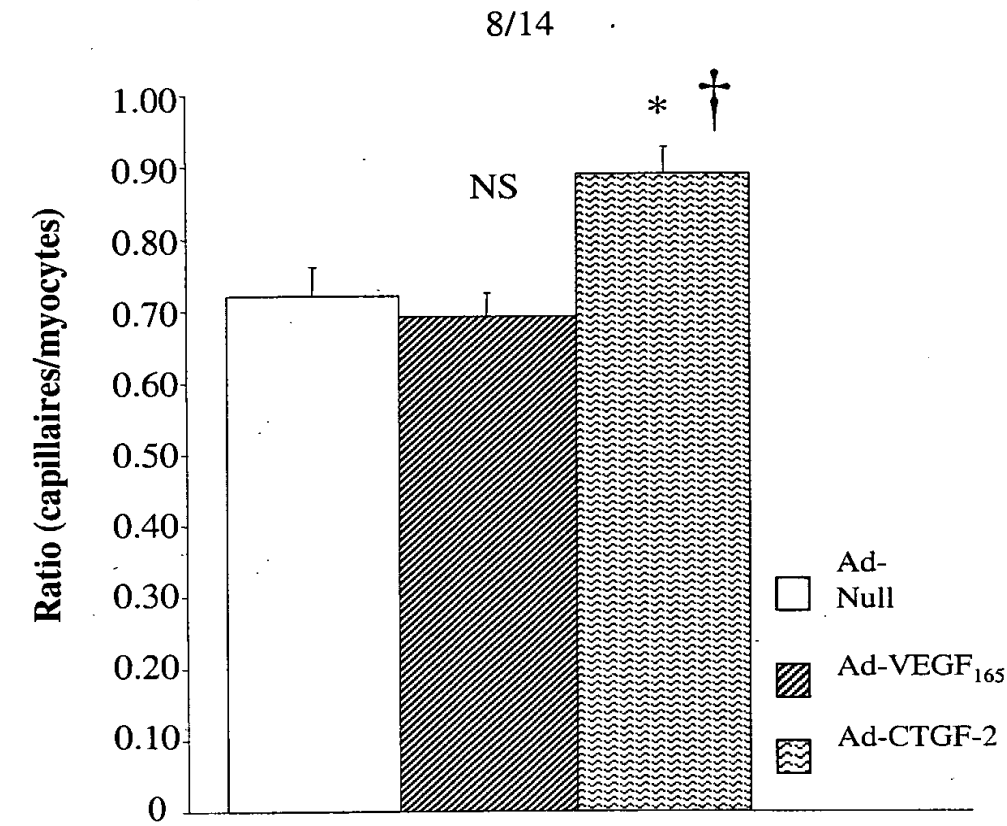


Figure 6

**Figure 7**

9/14

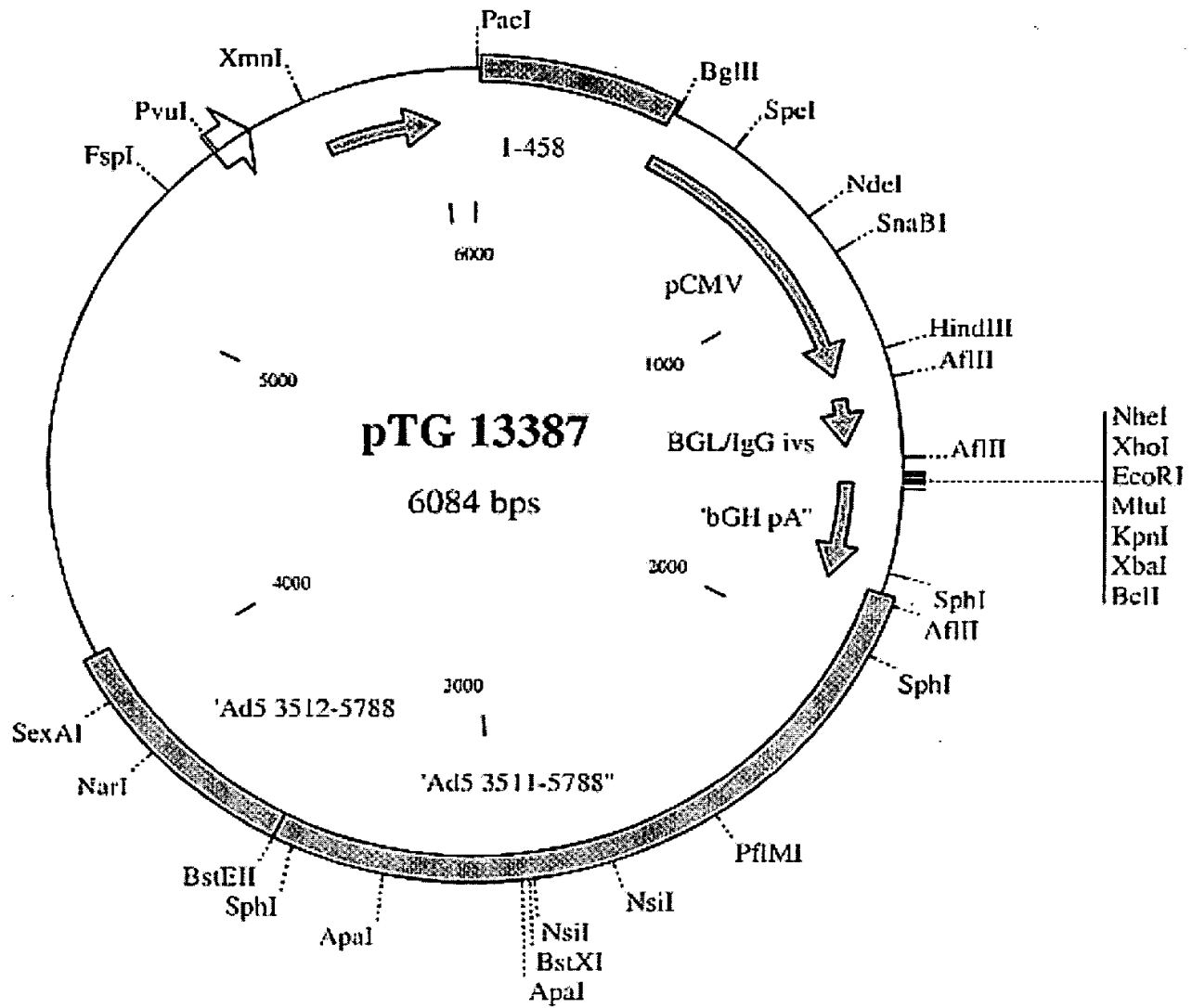


Figure 8

10/14

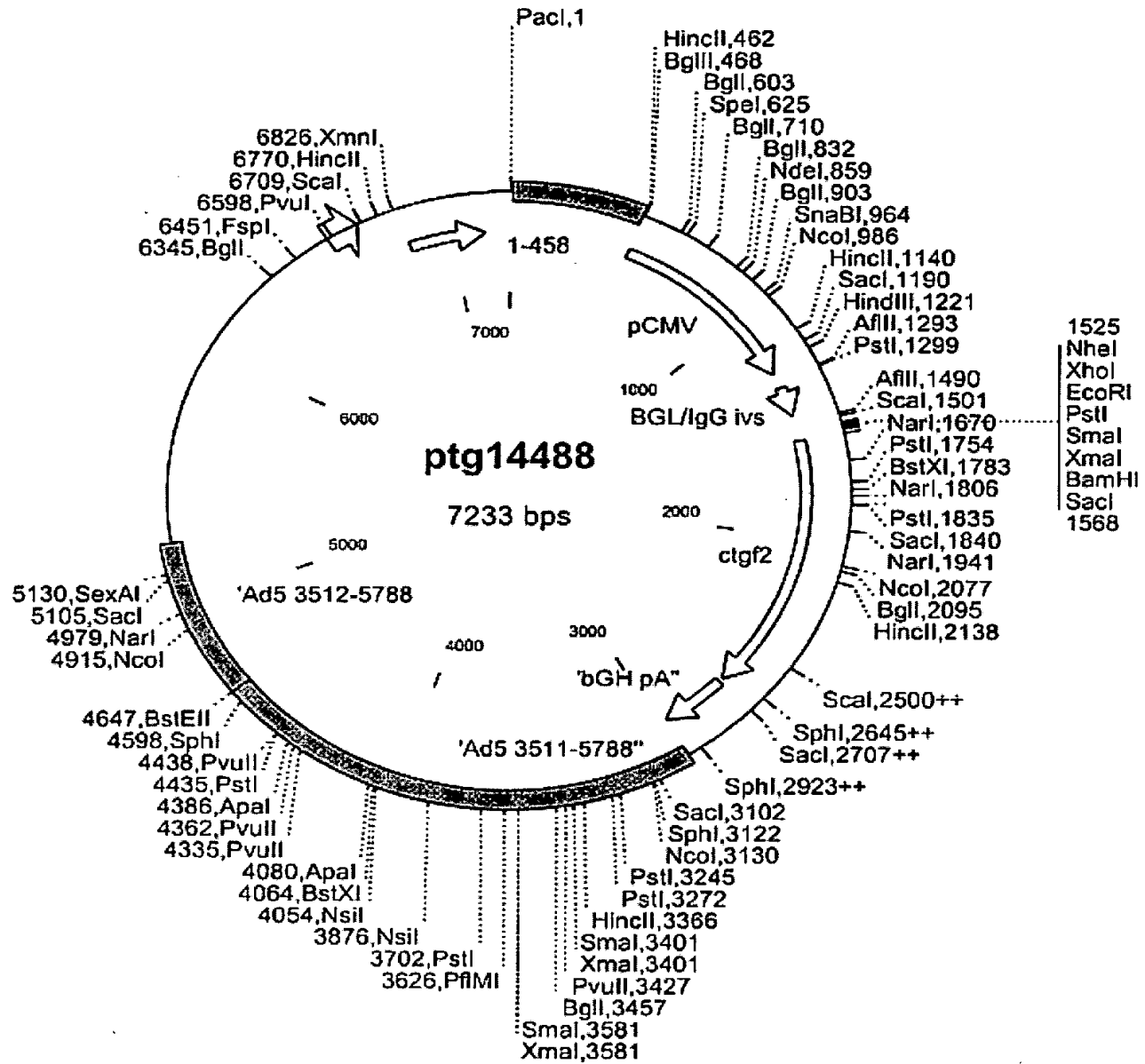


Figure 9

Figure 10
11/14

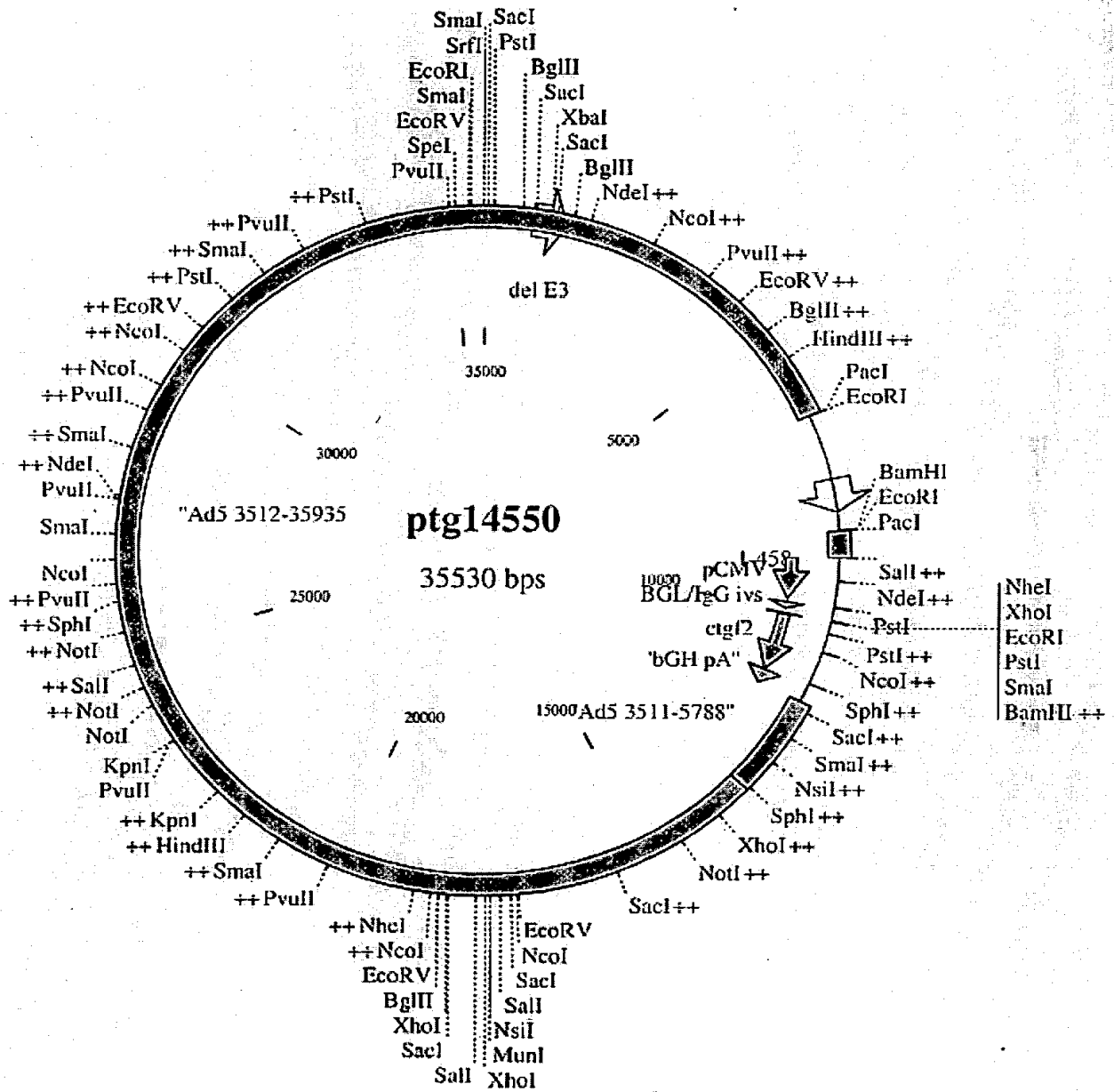


Figure 11A
12/14

ATGAGCTCCGGAATCGTCAGGAGCTCGCCTTAGTCGTACCCCTTCTCCACTTGACCAGG
M S S R I V R E L A L V V T L L H L T R
GTGGGGCTCTCCACCTGCCCGCTGACTGCCACTGCCCGCTGGAGGGCGCCCAAGTGC GCG
V G L S T C P A D C H C P L E A P K C A
CCGGGAGTCGGGCTGGTCCGGGACGGCTGCCGCTGTTGTAAGTCTGCGCCCAAGCAGCTC
P G V G L V R D G C G C C K V C A K Q L
AACGAGGACTGCAGAAAACGCGAGCCTGCCGACCACACCAAGGGCTGGAATGCAACTTC
N E D C R K T Q P C D H T K G L E C N F
GGCGCCAGCTCCACCGCTCTGAAGGGGATCTGCAGAGCTCAGTCAGAGGGCAGACCCCTGT
G A S S T A L K G I C R A Q S E G R P C
GAATATACTCCAGAAATCTACCAAACGGGGAAAGTTTCCAGCCCAACTGTAAACATCAG
E Y N S R I Y Q N G E S F Q P N C K H Q
TGCACATGTATTGGATGGCGCGGGGGCTTGCAATTCCTCTGTGTCCCCCAAGAACTATCT
C T C I G W R R G A C I P L C P Q E L S

09901910.070202

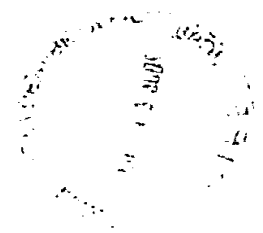


Figure 11B
13/14

CTCCCAACTTGGGCTGTCCCAACCCCTCGGCTGGTCAAAGTTACCGGGCAGTGCTGGGAG
L P N L G C P N P R L V K V T G Q C C E

GAGTGGTCTGTGACGAGGATAGTATCAAGGACCCCATGGAGGACCAGGACGGCCTCCTT
E W V C D E D S I K D P M E D Q D G L L

GGCAAGGGGCTGGGATTTCGATGCCTCCGAGGTGGAGTTGACGAGAAACAATGAATTGATT
G K G L G F D A S E V E L T R N E L I

GCAGTTGGAAGGAGCAGCTCACTGAAGCGGCTCCCTGTTTGGAAATGGAGCCCTCGCATC
A V G K G S S L K R L P V F G M E P R I

CTATACAACCCCTTTACAAGGCCAGAAATGTATTGTTCAAACAACCTTCATGGTCCCAGTGC
L Y N P L Q G Q K C I V Q T T S W S Q C

TCAAAGACCTGTGGAAGTGGTATCTCCACACGAGTTACCAATGACAAACCCTGAGTGCCGC
S K T C G T G I S T R V T N D N P E C R

CTTGTAAGAAAGAACCCGGATTGTGAGGTGCGGCTTGTGGACAGCCAGTGTAACAGCAGC
L V K E T R I C E V R P C G Q P V Y S S

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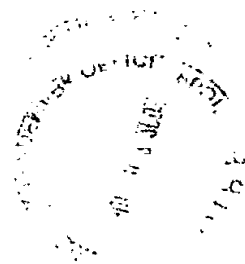


Figure 11C
14/14

CTGAAAAGGGCAAGAAATGCAGCAAGACCAAGAAATCCCCCGAACCAAGTCAGGTTTACT
L K K G K K C S K T K K S P E P V R F T

TACGCTGGATGTTTGAGTGTGAAGAAATACCGGCCCAAGTACTGCGGTTCTCCTGCGTGGAC
Y A G C L S V K K Y R P K Y C G S C V D

GGCCGATGCTGCACGCCCCAGCTGACCAAGGACTGTGAAGATGCGGTTCCCTGCGAAGAT
G R C C T P Q L T R T V K M R F P C E D

GGGAGAGACATTTTCCAAGAACGTCATGATGATCCAGTCCCTCCAAATGCAACTACACTGC
G E T F S K N V M M I Q S S K C N Y N C

CCGCATGCCAATGAAGCAGCGTTTCCCTTCTACAGGCTGTTCCTCAATGA
P H A N E A A F P F Y R L F Q .

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